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No Amendments are made in this response. Therefore, this listing of claims is provided for the convenience of the Examiner:

Listing of Claims:

(Original) A method for targeted gene repair, comprising
contacting a non-repaired target RNA with an RNA oligonucleotide complex comprising
a first oligonucleotide and a second oligonucleotide, said first oligonucleotide comprising a
sequence complementary to a repaired target RNA, wherein the RNA sequence of the first
oligonucleotide comprises an RNase H-resistant modification, and said second oligonucleotide
comprises an RNA sequence complementary to at least 6 nucleotides of the first oligonucleotide
at the site in the sequence of the first oligonucleotide which is not complementary to the nonrepaired target RNA; and

hybridizing said complex to said non-repaired target RNA in the presence of an RNase, wherein a repaired RNA is produced.

- (Original) The method of claim 1, wherein the repaired target RNA comprises a wild-type sequence.
- (Original) The method of claim 2, wherein the non-repaired target RNA comprises a
 mutation compared to said wild type sequence.
- (Original) The method of claim 3, wherein said mutation is a substitution, deletion or insertion of at least one base pair compared to said wild type sequence.
- 5. (Original) The method of claim 1, further comprising, preceding the steps of claim 1, contacting the non-repaired target RNA with a phosphorothoiate (PS) containing sequence comprising a deoxynucleotide with RNase H resistant flanking ends.
- (Original) The method of claim 1, wherein said RNase H-resistant modification is the addition of a 2-O-methyl moiety.

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- (Original) The method of claim 1, wherein said first oligonucleotide is at least 10 nucleotides in length.
- (Original) The method of claim 7, wherein said first oligonucleotide comprises about 33 nucleotides.
- (Original) The method of claim 1, wherein said second oligonucleotide comprises at least 7 nucleotides.
- (Original) The method of claim 9, wherein said second oligonucleotide comprises about 11 nucleotides.
- (Original) The method of claim 1, wherein said first oligonucleotide and said second oligonucleotide are annealed.
- (Original) The method of claim 1, wherein contacting said target RNA occurs within a
- 13. (Original) The method of claim 12, wherein said cell is in vitro, ex vivo or in vivo.
- 14. (Original) The method of claim 12, wherein said cell is a human cell.
- (Original) A method for treating or ameliorating a symptom of cystic fibrosis in a subject in need thereof, comprising

administering an RNA oligonucleotide complex directed to a non-repaired target RNA, said complex comprising a first oligonucleotide and a second oligonucleotide, said first oligonucleotide comprising a sequence complementary to a repaired target RNA, wherein the RNA sequence of the first oligonucleotide comprises an RNase H-resistant modification, and said second oligonucleotide comprises an RNA sequence complementary to at least 6 nucleotides

of the first oligonucleotide at the site on the sequence of the first oligonucleotide which is not complementary to the non-repaired target RNA; and

wherein administration produces a repaired targeted RNA, thereby treating or ameliorating symptom of cystic fibrosis.

- (Original) The method of claim 15, wherein the repaired target RNA comprises a wildtype sequence.
- (Original) The method of claim 16, wherein the non-repaired target RNA comprises a
 mutation compared to said wild type sequence.
- 18. (Original) The method of claim 17, wherein said mutation is a substitution, deletion or insertion of at least one base pair compared to said wild type sequence.
- 19. (Original) The method of claim 15, further comprising, preceding the steps of claim 15, administering a phosphorothoiate (PS) containing sequence comprising a deoxynucleotide with RNase H resistant flanking ends.
- (Original) The method of claim 15, wherein said RNase H-resistant modification is the addition of a 2-O-methyl moiety.
- (Original) The method of claim 15, wherein said first oligonucleotide is at least 10 nucleotides in length.
- (Original) The method of claim 21, wherein said first oligonucleotide comprises about 33 nucleotides.
- (Original) The method of claim 15, wherein said second oligonucleotide comprises at least 7 nucleotides.

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- (Original) The method of claim 23, wherein said second oligonucleotide comprises about
 11 nucleotides.
- (Original) The method of claim 15, wherein said first oligonucleotide and said second oligonucleotide are annealed.
- 26. (Original) An RNA oligonucleotide complex for modulating the expression or activity of a cystic fibrosis transmembrane conductance regulator (CFTR) gene product, the complex comprising a first oligonucleotide and a second oligonucleotide, said first oligonucleotide comprising the nucleic acid sequence of SEQ ID NO: 1 and said second oligonucleotide comprising the nucleic acid sequence of SEQ ID NO: 2, wherein said first and second oligonucleotide are annealed.

27-40. (Cancelled)